

#2
OIKE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/897,778

DATE: 07/20/2001
TIME: 14:08:00

Input Set : A:\455c16.app.txt
Output Set: N:\CRF3\07202001\I897778.raw

4 <110> APPLICANT: Wang, Tongtong
5 Marnerakis, Margarita
6 Fanger, Gary R.
7 Vedvick, Thomas S.
8 Carter, Darrick
9 Watanabe, Yoshihiro
10 Henderson, Robert A.
11 Peckham, David W.
12 Fanger, Neil

16 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
17 AND DIAGNOSIS OF LUNG CANCER

20 <130> FILE REFERENCE: 210121.455C16

C--> 22 <140> CURRENT APPLICATION NUMBER: US/09/897,778

23 <141> CURRENT FILING DATE: 2001-06-28

25 <160> NUMBER OF SEQ ID NOS: 467

27 <170> SOFTWARE: FastSEQ for Windows Version 4.0

29 <210> SEQ ID NO: 1

30 <211> LENGTH: 315

31 <212> TYPE: DNA

32 <213> ORGANISM: Homo sapiens

34 <220> FEATURE:

35 <221> NAME/KEY: misc_feature

36 <222> LOCATION: 236, 241

37 <223> OTHER INFORMATION: n = A,T,C or G

39 <400> SEQUENCE: 1

40 gcagagacag actggtggtt gaacctggag gtgccaaaaa agccagctgc gggcccagga 60
41 cagctgccgt gagactcccg atgtcacagg cagtctgtgt ggttacagcg cccctcagtg 120
42 ttcattctcca gcagagacaa cggaggaggc tcccaccagg acggttctca ttatttatat 180
W--> 43 gtaaatatgt ttgtaaactc atgtacagtt ttttttgggg gggaagcaat gggaanggta 240
W--> 44 naaattacaa atagaatcat ttgctgtaat ccttaaatgg caaacggtca ggccacgtga 300
45 aaaaaaaaaa aaaaaa 315

47 <210> SEQ ID NO: 2

48 <211> LENGTH: 380

49 <212> TYPE: DNA

50 <213> ORGANISM: Homo sapiens

52 <400> SEQUENCE: 2

53 atttaggctt aagattttgt ttacccttgt tactaaggag caaattagta ttaaagtata 60
54 atatatataa acaaatataa aaagttttga gtggttcagc ttttttatat tttttaatgg 120
55 cataactttt aacaacactg ctctgtaatg ggttgaactg tggtaactcag actgagataa 180
56 ctgaaatgag tggatgtata gtgttattgc ataattatcc cactatgaag caaagggact 240
57 ggataaattc ccagctctaga ttattagcct ttgttaacca tcaagcacct agaagaagaa 300
58 ttattggaaa ttttgtcctc tgtaactggc actttggggg gtgacttatc ttttgccctt 360
59 gtaaaaaaaaaa aaaaaaaaaa 380

61 <210> SEQ ID NO: 3

62 <211> LENGTH: 346

63 <212> TYPE: DNA

64 <213> ORGANISM: Homo sapiens

ENTERED
see page 5

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66 <220> FEATURE:
67 <221> NAME/KEY: misc_feature
68 <222> LOCATION: 316, 317, 318, 322, 323, 326, 329, 330, 331, 336, 337, 339,
69     340, 342, 343
70 <223> OTHER INFORMATION: n = A,T,C or G
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73 ttgtaagtat acaatttttag aaaggattaa atgttattga tcattttact gaatactgca 60
74 caccctcacc atacaccatc cactttccaa taacatttaa tcctttctaa aattgtaagt 120
75 atacaattgt actttctttg gattttcata acaaataac catagactgt taattttatt 180
76 gaagtttcct taatggaatg agtcattttt gtcttggtct ttgagggtta ctttgcttt 240
77 gacttccaac aatttgatca tatagtgttg agctgtggaa atctttaagt ttattctata 300
W--> 78 gcaataattt ctattnnnag annccnggnn naaaannann annaaa 346
80 <210> SEQ ID NO: 4
81 <211> LENGTH: 372
82 <212> TYPE: DNA
83 <213> ORGANISM: Homo sapiens
85 <220> FEATURE:
86 <221> NAME/KEY: misc_feature
87 <222> LOCATION: 297, 306, 332
88 <223> OTHER INFORMATION: n = A,T,C or G
90 <400> SEQUENCE: 4
91 actagtctca ttactccaga attatgctct tgtacctgtg tggctgggtt tcttagtctg 60
92 tggtttggtt tggttttttg aactgggatg taggggtggtt cacagttcta atgtaagcac 120
93 tctcttctcc aagttgtgct ttgtggggac aatcattctt tgaacattag agaggaaggc 180
94 agttcaagct gttgaaaaga ctattgctta tttttgtttt taaagacctt cttgacgtca 240
W--> 95 tgtggacagt gcacgtgcct tacgctacat cttgttttct aggaagaagg ggatgcnngg 300
W--> 96 aaggantggg tgctttgtga tggataaaac guctaaataa cacaccttta cattttgaaa 360
97 aaaacaaaac aa 372
99 <210> SEQ ID NO: 5
100 <211> LENGTH: 698
101 <212> TYPE: DNA
102 <213> ORGANISM: Homo sapiens
104 <220> FEATURE:
105 <221> NAME/KEY: misc_feature
106 <222> LOCATION: 8, 345, 422, 430, 433, 436, 438, 472, 481, 486, 515, 521,
107     536, 549, 553, 556, 557, 559, 568, 593, 597, 605, 611, 613,
108     616, 618, 620, 628, 630, 632, 634, 635, 639, 643, 647, 648,
109     649, 652, 654, 658, 664, 690
110 <223> OTHER INFORMATION: n = A,T,C or G
112 <400> SEQUENCE: 5
W--> 113 actagtanga tagaaacact gtgtcccgag agtaaggaga gaagctacta ttgattagag 60
114 cctaaccag gtttaactgca agaagaggcg ggatactttc agctttccat gtaactgtat 120
115 gcataaagcc aatgtagtcc agttttctaag atcatgttcc aagctaactg aatcccactt 180
116 caatacacac tcatgaactc ctgatggaac aataacaggc ccaagcctgt ggtatgatgt 240
117 gcacacttgc tagactcaga aaaaatacta ctctcataaa tgggtgggag tattttgggt 300
W--> 118 gacaacctac tttgcttggc tgagtgaagg aatgatattc atatnttcat ttattccatg 360
119 gacatttagt tagtgctttt tatataccag gcatgatgct gagtgaactc cttgtgtata 420
W--> 120 tntccaaatn ttngtnngt cgctgcacat atctgaaatc ctatattaag antttccaa 480
W--> 121 natgangtcc ctggtttttc cagccactt gatcngtcaa ngatctcacc tctgtntgtc 540

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W--> 122 ctaaacacnt ctnctnnaag gttagacnng acctctcttc tcccttcccg aanaathnaag 600
W--> 123 tgtgngaaga nancncncn cccccctnch tcnncctng ccngetnnnc cncntgtngg 660
W--> 124 gggngcgcgc cccgcggggg gaccccccn ttttcccc 698
126 <210> SEQ ID NO: 6
127 <211> LENGTH: 740
128 <212> TYPE: DNA
129 <213> ORGANISM: Homo sapiens
131 <220> FEATURE:
132 <221> NAME/KEY: misc_feature
133 <222> LOCATION: 82, 406, 426, 434, 462, 536, 551, 558, 563, 567, 582, 584,
134 592, 638, 651, 660, 664, 673, 675, 697, 706, 711, 715, 716,
135 717, 723, 724, 725, 733
136 <223> OTHER INFORMATION: n = A,T,C or G
138 <400> SEQUENCE: 6
139 actagtcaaa aatgctaaaa taatttgga gaaaatattt tttaagtagt gttatagttt 60
W--> 140 catgtttatc ttttattatg tnttgtgaag ttgtgtcttt tcaactaatta cctatactat 120
141 gccaatattt ccttatactc atccataaca ttatactac atttgtaaga gaatatgcac 180
142 gtgaaactta acactttata aggtaaaaat gaggtttcca agatttaata atctgatcaa 240
143 gttcttggtta tttccaaata gaatggactt ggtctgttaa ggggctaagg gagaagaaga 300
144 agataagggtt aaaagttgtt aatgaccaa cattctaaaa gaaatgcaa aaaaaattta 360
W--> 145 ttttcaagcc ttcgaactat ttaaggaaag caaaatcatt tcttanatgc atatcatttg 420
W--> 146 tgagantttc tcantaatat cctgaatcat tcatttcagc tnaggcttca tgttgatcg 480
W--> 147 atatgtcatc tagggaaagt ctatttcag gtccaaacct gttgcatag ttggttaggc 540
W--> 148 tttcctttaa ntgtgaanta ttnacangaa attttctctt tnanagttct tnataggggt 600
W--> 149 aggggtgtgg gaaaagcttc taacaatctg tagtgttncg tgttatctgt ncagaaccan 660
W--> 150 aatgacggat cgnangaagg actgggtcta tttacangaa cgaatnatct ngttnnntgt 720
W--> 151 gtnnncaact cchgggagcc 740
153 <210> SEQ ID NO: 7
154 <211> LENGTH: 670
155 <212> TYPE: DNA
156 <213> ORGANISM: Homo sapiens
158 <220> FEATURE:
159 <221> NAME/KEY: misc_feature
160 <222> LOCATION: 265, 268, 457, 470, 485, 546, 553, 566, 590, 596, 613, 624,
161 639, 653, 659, 661
162 <223> OTHER INFORMATION: n = A,T,C or G
164 <400> SEQUENCE: 7
165 gctggggagc tcggcatggc ggtccccgct gcagccatgg ggcctcggc gttgggccag 60
166 agcgcccccg gctcgatggc cccgtggtgc tcagtgaaga gcggccccgc gcgctacgtg 120
167 cttgggatgc aggagctgtt ccggggccac agcaagaccg cgagttctcg gcgcacagcg 180
168 ccaagggtgca ctcggtggcc tggagtggcg acgggcgtcg cctacctcgg ggtcttcgac 240
W--> 169 aagacgccac gtcttcttgc tgganaanga ccgttgggtc aagaaaacaa ttatcgggga 300
170 catggggata gtgtggacca ctttgttggc atccaagtaa tctgaccta tttgttacgg 360
171 cgtctggaga taaaaccatt cgcactctgg atgtgaggac taaaaaatgc attgccactg 420
W--> 172 tgaacactaa aggggagaac attaatatct gctggantcc tgatgggcan accattgtcg 480
W--> 173 tagcnacaag gatgatgtgg tgactttatt gatgccaaga aaccccgttc caaagcaaaa 540
W--> 174 aaacanttcc aanttcgaag tcacchaaat ctcttggaag aatgaacatn aatatnttct 600
W--> 175 tcttgacaat ggccttggg tgtntecat cctcagctnc cccaaaactg aancctgtnc 660
W--> 176 natccacccc 670

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178 <210> SEQ ID NO: 8
179 <211> LENGTH: 689
180 <212> TYPE: DNA
181 <213> ORGANISM: Homo sapiens
183 <220> FEATURE:
184 <221> NAME/KEY: misc_feature
185 <222> LOCATION: 253, 335, 410, 428, 448, 458, 466, 479, 480, 482, 483, 485,
186      488, 491, 492, 495, 499, 500, 502, 503, 512, 516, 524, 525,
187      526, 527, 530, 540, 546, 550, 581, 593, 594, 601, 606, 609,
188      610, 620, 621, 622, 628, 641, 646, 656, 673
189 <223> OTHER INFORMATION: n = A,T,C or G
191 <400> SEQUENCE: 8
192 actagtatct aggaatgaac agtaaaagag gagcagttgg ctacttgatt acaacagagt 60
193 aaatgaagta ctggatttgg gaaaacctgg ttttattaga acatatggaa tgaaagccta 120
194 cacctagcat tgcctactta gcccctgaa ttaacagagc ccaattgaga caaaccctcg 180
195 gcaacaggaa attcaaggga gaaaaagtaa gcaacttggg ctaggatgag ctgactccct 240
W--> 196 tagagcaaag ganagacagc cccattacc aaataccatt ttgctctggg gcttctgcag 300
W--> 197 ctggcagtgt tcctgcccc gcatggcacc ttatngtttt gatagcaact tcgttgaatt 360
W--> 198 ttcaccaact tattacttga aattataaja tagcctgtgc gtttgcgtg tccaggctgt 420
W--> 199 gatatatntt cctagtgttt tgacttttaa aataaainag gtttannttt cccccccnn 480
W--> 200 cnnctctncc nntcnctcnn cnncccccc cncctngtcc tccnnnttn gggggggccn 540
W--> 201 cccccnegg ggacccccct tgggtccctt agtggagggt natgggcctt ggnnttatcc 600
W--> 202 nggccttann ttccccctn nnaaatgntt cccctccca ntcccnccac ctcaanccgg 660
W--> 203 aagcctaagt ttntaccctg ggggtcccc 689
205 <210> SEQ ID NO: 9
206 <211> LENGTH: 674
207 <212> TYPE: DNA
208 <213> ORGANISM: Homo sapiens
210 <220> FEATURE:
211 <221> NAME/KEY: misc_feature
212 <222> LOCATION: 602, 632, 639, 668
213 <223> OTHER INFORMATION: n = A,T,C or G
215 <400> SEQUENCE: 9
216 gtccactctc ctttgagtgt actgtcttac tgtgcactct gtttttcaac tttctagata 60
217 taaaaaatgc ttgttctata gtggagtaag agctcacaca cccaaggcag caagataact 120
218 gaaaaaagcg aggccttttt gccaccttgg taaaggccag ttcactgcta tagaactgct 180
219 ataagcctga agggaagtag ctatgagact ttccattttt cttagtcttc ccaataggct 240
220 ccttcattga aaaaggcttc ctgtaataat ttccacctaa tgaattagca gtgtgattat 300
221 ttctgaaata agagacaaat tgggcccag agtcttcttg tgatttaaaa taaacaaccc 360
222 aaagttttgt ttggtcttca ccaaaggaca tactctaggg ggtatgttgt tgaagacatt 420
223 caaaaacatt agctgttctg tctttcaatt tcaagttatt ttggagactg cctccatgtg 480
224 agttaattac tttgctctgg aactagcatt attgtcatta tcatcacatt ctgtcatcat 540
225 catctgaata atattgtgga tttccccctc tgggttgcac ttcttttgac tctctggga 600
W--> 226 anaaatgtca aaaaaaagg tcgatctact cngcaaggnc catctaata ctgcgctgga 660
W--> 227 aggaccnct gcc 674
229 <210> SEQ ID NO: 10
230 <211> LENGTH: 346
231 <212> TYPE: DNA
232 <213> ORGANISM: Homo sapiens

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234 <220> FEATURE:
235 <221> NAME/KEY: misc_feature
236 <222> LOCATION: 320, 321, 322, 325, 326, 328, 329, 330, 332, 333, 334, 335,
237      342
238 <223> OTHER INFORMATION: n = A,T,C or G
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242 ttctgtctgt aacaaaaatg tactttatag agatggagga aaaggtctaa tactacatag 120
243 ccttaagtgt ttctgtcatt gttcaagtgt attttctgta acagaaacat atttggaaatg 180
244 tttttctttt ccccttataa attgtaattc ctgaaatact gctgctttaa aaagtccac 240
245 tgtcagatta tattatctaa caattgaata ttgtaaatat agttgtctta cctctcaata 300
W--> 246 aaaggtact tttctattan nnagngnnn gnnnnataaa anaaaa 346
248 <210> SEQ ID NO: 11
249 <211> LENGTH: 602
250 <212> TYPE: DNA
251 <213> ORGANISM: Homo sapiens
253 <400> SEQUENCE: 11
254 actagtataaa agcagcattg ccaaataatc cctaattttc cactaaaaat ataatgaaat 60
255 gatgttaagc tttttgaaaa gttaggta aacctactgt tgtagatta atgtatttgt 120
256 tgcttccctt tatctggaat gtgcattag cttttttatt ttaacctctt ttaattctta 180
257 ttcaattcca tgacttaagg ttggagagct aaacactggg atttttggaat aacagactga 240
258 cagttttgca taattataat cggcattgta catagaaagg atatggctac cttttgttaa 300
259 atctgcactt tctaaatata aaaaaaggga aatgaagtta taaatcaatt ttgtataat 360
260 ctgtttgaaa catgagtttt atttgcttaa tattagggct ttgccctttt tctgtaagtc 420
261 tcttgggata ctgtgtagaa ctgttctcat taaacaccaa acagtttaagt ccattctctg 480
262 gtactagcta caaattcggg ttcatattct acttaacaat taaataaac tgaaatattt 540
263 ctagatggtc tactttctgt catataaaaa caaaacttga tttccaaaaa aaaaaaaaaa 600
264 aa 602
266 <210> SEQ ID NO: 12
267 <211> LENGTH: 685
268 <212> TYPE: DNA
269 <213> ORGANISM: Homo sapiens
271 <220> FEATURE:
272 <221> NAME/KEY: misc_feature
273 <222> LOCATION: 170, 279, 318, 321, 322, 422, 450, 453, 459, 467, 468, 470,
274      473, 475, 482, 485, 486, 491, 498, 503, 506, 509, 522, 526,
275      527, 528, 538, 542, 544, 551, 567, 568, 569, 574, 576, 582,
276      587, 588, 589, 590, 592, 593, 598, 599, 603, 605, 608
277 <223> OTHER INFORMATION: n = A,T,C or G
279 <221> NAME/KEY: misc_feature
280 <222> LOCATION: 633, 634, 635, 644, 646, 648, 651, 655, 660, 662, 663, 672,
281      674, 675, 682, 683
282 <223> OTHER INFORMATION: n = A,T,C or G
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286 attatcatgg tattgatgga cctaagaaaa taaaaattag actaagcccc caaataagct 120
W--> 287 gcatgcattt gtaacatgat tagtagattt gaatatatag atgtagtatt ttgggtatct 180
288 aggtgtttta tcattatgta aaggaattaa agtaaaggag ttgttagttg tttttattaa 240
W--> 289 atatgcatat agtagagtgc aaaaatatag caaaaatana aactaaaggt agaaaagcat 300

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Use of n and / or Xaa has been detected in the
Sequence Listing. Review the Sequence Listing
to ensure a corresponding explanation is present
in the <220> to <223> fields of each sequence
using n or Xaa.

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L:22 M:270 C: Current Application Number differs, Wrong Format
L:43 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:148 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:150 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:199 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:289 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:293 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:318 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13

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L:320 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13

L:321 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13

L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14